

=> File .Biotech

=> s (memapsin 2 or beta secretase or beta amyloid precursor protein or APP)  
L1 630677 (MEMAPSIN 2 OR BETA SECRETASE OR BETA AMYLOID PRECURSOR PROTEIN  
OR APP)

=> s (l1 or memapsin 2) and (OM 99 or OM-99 or dipeptide isostere or dipeptide or  
isostere) and Alzheimer?  
L2 135 (L1 OR MEMAPSIN 2) AND (OM 99 OR OM-99 OR DIPEPTIDE ISOSTERE OR  
DIPEPTIDE OR ISOSTERE) AND ALZHEIMER?

=> s l2 and (computer program# or software program#)  
L3 22 L2 AND (COMPUTER PROGRAM# OR SOFTWARE PROGRAM#)

=> s l3 and (recombinant?)  
L4 22 L3 AND (RECOMBINANT?)

=> s l4 and (treat? or therapeut? or diagnos? or prevent?)  
5 FILES SEARCHED...  
L5 22 L4 AND (TREAT? OR THERAPEUT? OR DIAGNOS? OR PREVENT?)

=> s l5 and (inhibit?)  
L6 22 L5 AND (INHIBIT?)

=> s Tang J?/au; s Hong L/au; s Ghosh A/au; s Koelsch G/au  
L7 8203 TANG J?/AU

L8 417 HONG L/AU

L9 1933 GHOSH A/AU

L10 100 KOELSCH G/AU

=> s l6 and (l7 or l8 or l9 or l10)  
L11 7 L6 AND (L7 OR L8 OR L9 OR L10)

=> d l11 1-7 bib ab

L11 ANSWER 1 OF 7 CAPLUS COPYRIGHT 2003 ACS on STN  
AN 2001:12489 CAPLUS  
DN 134:80832  
TI Inhibitors of memapsin 2 and use thereof  
IN Tang, Jordan J. N.; Hong, Ling; Ghosh, Arun K.  
PA Oklahoma Medical Research Foundation, USA; The Board of Trustees of the  
University of Illinois  
SO PCT Int. Appl., 86 pp.  
CODEN: PIXXD2  
DT Patent  
LA English  
FAN.CNT 3

	PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
PI	WO 2001000665	A2	20010104	WO 2000-US17742	20000627
	WO 2001000665	A3	20010927		
	WO 2001000665	C2	20020725		
	W:	AE, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CR, CU, CZ, DE, DK, DM, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, UZ, VN, YU, ZA, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM			
	RW:	GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY,			

DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF, BJ,  
 CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG  
 EP 1194449 A2 20020410 EP 2000-943236 20000627  
 R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT,  
 IE, SI, LT, LV, FI, RO  
 JP 2003506322 T2 20030218 JP 2001-507071 20000627  
 US 6545127 B1 20030408 US 2000-604608 20000627  
 US 2002049303 A1 20020425 US 2001-796264 20010228  
 US 2002164760 A1 20021107 US 2001-795903 20010228  
 US 2002115600 A1 20020822 US 2001-845226 20010430  
 PRAI US 1999-141363P P 19990628  
 US 1999-168060P P 19991130  
 US 2000-177836P P 20000125  
 US 2000-178368P P 20000127  
 US 2000-210292P P 20000608  
 US 2000-603713 A3 20000627  
 US 2000-604608 A3 20000627  
 WO 2000-US17742 W 20000627  
 AB Methods for the prodn. of purified, catalytically active,  
 recombinant memapsin 2 have been developed.  
 The substrate and subsite specificity of the catalytically active enzyme  
 have been detd. The substrate and subsite specificity information was  
 used to design substrate analogs of the natural memapsin  
 2 substrate that can inhibit the function of  
 memapsin 2. The substrate analogs are based on peptide  
 sequences, shown to be related to the natural peptide substrates for  
 memapsin 2. The substrate analogs contain at least one  
 analog of an amide bond which is not capable of being cleaved by  
 memapsin 2. Processes for the synthesis of two  
 substrate analogs including isosteres at the sites of the crit. amino acid  
 residues were developed and the substrate analogs, OM99-1 and OM99-2,  
 were synthesized. OM99-2 is based on an octapeptide Glu-Val-Asn-Leu-Ala-  
 Ala-Glu-Phe (SEQ ID NO:28) with the Leu-Ala peptide bond substituted by a  
 transition-state isostere hydroxyethylene group (Figure 1). The  
 inhibition const. of OM99-2 is 1.6 x 10-9 M against  
 recombinant pro-memapsin 2. Crystallog. of  
 memapsin 2 bound to this inhibitor was used to  
 det. the three dimensional structure of the protein, as well as the  
 importance of the various residues in binding. This information can be  
 used by those skilled in the art to design new inhibitors, using  
 com. available software programs and techniques  
 familiar to those in org. chem. and enzymol., to design new  
 inhibitors to memapsin 2, useful in  
 diagnostics and for the treatment and/or  
 prevention of Alzheimer's disease.

L11 ANSWER 2 OF 7 CAPLUS COPYRIGHT 2003 ACS on STN  
 AN 2001:12487 CAPLUS  
 DN 134:68049  
 TI Catalytically active recombinant memapsin 2,  
 3D crystal structure based inhibitor design, synthesis, and  
 screening, for Alzheimer's disease treatment  
 IN Tang, Jordan J. N.; Lin, Xinli; Koelsch, Gerald  
 PA Oklahoma Medical Research Foundation, USA  
 SO PCT Int. Appl., 87 pp.  
 CODEN: PIXXD2  
 DT Patent  
 LA English  
 FAN.CNT 3

	PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
PI	WO 2001000663	A2	20010104	WO 2000-US17661	20000627
	WO 2001000663	A3	20011004		
	W: AE, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CR, CU, CZ, DE, DK, DM, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL,				

IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, UZ, VN, YU, ZA, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM  
RW: GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG  
EP 1196609 A2 20020417 EP 2000-943208 20000627  
R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT, IE, SI, LT, LV, FI, RO  
JP 2003503072 T2 20030128 JP 2001-507069 20000627  
US 6545127 B1 20030408 US 2000-604608 20000627  
US 2002049303 A1 20020425 US 2001-796264 20010228  
US 2002164760 A1 20021107 US 2001-795903 20010228  
US 2002115600 A1 20020822 US 2001-845226 20010430  
PRAI US 1999-141363P P 19990628  
US 1999-168060P P 19991130  
US 2000-177836P P 20000125  
US 2000-178368P P 20000127  
US 2000-210292P P 20000608  
US 2000-603713 A3 20000627  
US 2000-604608 A3 20000627  
WO 2000-US17661 W 20000627  
AB A method for producing catalytically active **recombinant memapsin 2** comprising expression in a bacteria and refolding the **recombinant memapsin 2** under conditions which dissociate and then slowly refold the enzyme into a catalytically active form is disclosed. A method of isolating **inhibitors** of cleavage by **memapsin 2** comprising adding to one or more potential **inhibitors** of catalytically active **recombinant memapsin 2**, and a substrate for **memapsin 2**, and screening for decreased cleavage of the substrate by the **inhibitors**, wherein the **inhibitors** are in a library of small synthetic mols., like proteins and peptides. Alternatively, the **inhibitors** are oligonucleotides preventing or decreasing expression of catalytically active **memapsin 2**. A method for designing or obtaining **inhibitors** of catalytically active **memapsin 2** comprising modeling an **inhibitor** based on the crystn. coordinates of **memapsin 2** or parameters. A database comprising binding properties and chem. structures of compds. designed or screened by modeling an **inhibitor** based on the crystn. coordinates of **memapsin 2** or parameters is claimed. A method of **treating or preventing Alzheimer's disease** comprising administering to a patient in need thereof an **inhibitor** of **memapsin 2** which binds to the active site of the **memapsin 2** defined by the presence of two catalytic aspartic residues and substrate binding cleft, is also claimed. The cDNAs of two new human membrane-assocd. aspartic proteases, **memapsin 1** and **memapsin 2**, have been cloned and sequenced. The substrate and subsite specificity of the catalytically active enzyme have been detd. The substrate and subsite specificity information was used to design substrate analogs of the natural **memapsin 2** substrate that can inhibit the function of **memapsin 2**. The substrate analogs are based on peptide sequences, shown to be related to the natural peptide substrates for **memapsin 2**. The substrate analogs contain at least one analog of an amide bond which is not capable of being cleaved by **memapsin 2**. Processes for the synthesis of two substrate analogs including isosteres at the sites of the crit. amino acid residues were developed and the substrate analogs, OMR99-1 and OM99-2, were synthesized. OM99-2 is based on an octapeptide Glu-Val-Asn-Leu-Ala-Ala-Glu-Phe (SEQ ID NO:28) with the Leu-Ala peptide bond substituted by a transition-state **isostere** hydroxyethylene group (Fig. 1). The **inhibition** const. of OM99-2 is 1.6 x 10<sup>9</sup> M

against recombinant pro-memapsin 2. Crystallog. of memapsin 2 bound to this inhibitor was used to det. the three dimensional structure of the protein, as well as the importance of the various residues in binding. This information can be used to design new inhibitors, using com. available software programs and techniques familiar to those in org. chem. and enzymol., to design new inhibitors to memapsin 2, useful in diagnostics and for the treatment and/or prevention of Alzheimer's disease.

L11 ANSWER 3 OF 7 USPATFULL on STN  
AN 2003:134541 USPATFULL  
TI Inhibitors of memapsin 2 and use thereof  
IN Tang, Jordan J. N., Edmond, OK, UNITED STATES  
Koelsch, Gerald, Oklahoma City, OK, UNITED STATES  
Ghosh, Arun K., River Forest, IL, UNITED STATES  
PA Oklahoma Medical Research Foundation, Oklahoma City, OK (U.S. corporation)  
PI US 2003092629 A1 20030515  
AI US 2001-32818 A1 20011228 (10)  
PRAI US 2001-275756P 20010314 (60)  
US 2000-258705P 20001228 (60)  
DT Utility  
FS APPLICATION  
LREP HAMILTON, BROOK, SMITH & REYNOLDS, P.C., 530 VIRGINIA ROAD, P.O. BOX 9133, CONCORD, MA, 01742-9133  
CLMN Number of Claims: 24  
ECL Exemplary Claim: 1  
DRWN 9 Drawing Page(s)  
LN.CNT 2203  
CAS INDEXING IS AVAILABLE FOR THIS PATENT.  
AB Methods for the production of purified, catalytically active, recombinant memapsin 2 have been developed-, The substrate and subsite specificity of the catalytically active enzyme have been determined by a method which determines the initial hydrolysis rate of the substrates by using MALDI-TOF/MS. Alternatively, the subsite specificity of memapsin can be determined by probing a library of inhibitors with memapsin 2 and subsequently detecting the bound memapsin 2 with an antibody raised to memapsin 2 and an alkaline phosphatase conjugated secondary antibody. The substrate and subsite specificity information was used to design substrate analogs of the natural memapsin 2 substrate that can inhibit the function of memapsin 2. The substrate analogs are based on peptide sequences, shown to be related to the natural peptide substrates for memapsin 2. The substrate analogs contain at least one analog of an amide bond which is not capable of being cleaved by memapsin 2. Processes for the synthesis of substrate analogues including isosteres at the sites of the critical amino acid residues were developed and the more than seventy substrate analogues were synthesized, among which MMI-005, MMI-012, MMI-017, MMI-018, MMI-025, MMI-026, MMI-037, MMI-039, MMI-040, MMI-066, MMI-070, and MMI-071 have inhibition constants in the range of 1.4-61.4.times.10.sup.-9 M against recombinant pro-memapsin 2. These inhibitors are useful in diagnostics and for the treatment and/or prevention of Alzheimer's disease.

L11 ANSWER 4 OF 7 USPATFULL on STN  
AN 2003:96167 USPATFULL  
TI Catalytically active recombinant memapsin and methods of use thereof  
IN Tang, Jordan J. N., Edmond, OK, United States  
Lin, Xinli, Edmond, OK, United States

Koelsch, Gerald, Oklahoma City, OK, United States  
Hong, Lin, Oklahoma City, OK, United States  
PA Oklahoma Medical Research Foundation, Oklahoma City, OK, United States  
(U.S. corporation)  
PI US 6545127 B1 20030408  
AI US 2000-604608 20000627 (9)  
PRAI US 1999-141363P 19990628 (60)  
US 1999-168060P 19991130 (60)  
US 2000-177836P 20000125 (60)  
US 2000-178368P 20000127 (60)  
US 2000-210292P 20000608 (60)  
DT Utility  
FS GRANTED  
EXNAM Primary Examiner: Borin, Michael; Assistant Examiner: Zhou, Shuba  
LREP Hamilton, Brook, Smith & Reynolds, P.C.  
CLMN Number of Claims: 18  
ECL Exemplary Claim: 7  
DRWN 21 Drawing Figure(s); 12 Drawing Page(s)  
LN.CNT 2563  
CAS INDEXING IS AVAILABLE FOR THIS PATENT.  
AB Methods for the production of purified, catalytically active, recombinant memapsin 2 have been developed. The substrate and subsite specificity of the catalytically active enzyme have been determined. The substrate and subsite specificity information was used to design substrate analogs of the natural memapsin 2 substrate that can inhibit the function of memapsin 2. The substrate analogs are based on peptide sequences, shown to be related to the natural peptide substrates for memapsin 2. The substrate analogs contain at least one analog of an amide bond which is not capable of being cleaved by memapsin 2. Processes for the synthesis of two substrate analogues including isosteres at the sites of the critical amino acid residues were developed and the substrate analogues, OMR99-1 and OM99-2, were synthesized. OM99-2 is based on an octapeptide Glu-Val-Asn-Leu-Ala-Ala-Glu-Phe (SEQ ID NO:28) with the Leu-Ala peptide bond substituted by a transition-state isostere hydroxyethylene group (FIG. 1). The inhibition constant of OM99-2 is 1.6.times.10.sup.-9M against recombinant pro-memapsin 2. Crystallography of memapsin 2 bound to this inhibitor was used to determine the three dimensional structure of the protein, as well as the importance of the various residues in binding. This information can be used by those skilled in the art to design new inhibitors, using commercially available software programs and techniques familiar to those in organic chemistry and enzymology, to design new inhibitors to memapsin 2, useful in diagnostics and for the treatment and/or prevention of Alzheimer's disease.  
L11 ANSWER 5 OF 7 USPATFULL on STN  
AN 2002:294717 USPATFULL  
TI Catalytically active recombinant memapsin and methods of use thereof  
IN Lin, Xinli, Edmond, OK, UNITED STATES  
Koelsch, Gerald, Oklahoma City, OK, UNITED STATES  
Tang, Jordan J.N., Edmond, OK, UNITED STATES  
PA Oklahoma Medical Research Foundation  
PI US 2002164760 A1 20021107  
AI US 2001-795903 A1 20010228 (9)  
RLI Division of Ser. No. US 2000-604608, filed on 27 Jun 2000, PENDING  
PRAI US 1999-141363P 19990628 (60)  
US 1999-168060P 19991130 (60)  
US 2000-177836P 20000125 (60)  
US 2000-178368P 20000127 (60)  
US 2000-210292P 20000608 (60)

DT Utility  
FS APPLICATION  
LREP PATREA L. PABST, HOLLAND & KNIGHT LLP, SUITE 2000, ONE ATLANTIC CENTER,  
1201 WEST PEACHTREE STREET, N.E., ATLANTA, GA, 30309-3400  
CLMN Number of Claims: 33  
ECL Exemplary Claim: 1  
DRWN 12 Drawing Page(s)  
LN.CNT 2440

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB Methods for the production of purified, catalytically active, recombinant **memapsin 2** have been developed. The substrate and subsite specificity of the catalytically active enzyme have been determined. The substrate and subsite specificity information was used to design substrate analogs of the natural **memapsin 2** substrate that can **inhibit** the function of **memapsin 2**. The substrate analogs are based on peptide sequences, shown to be related to the natural peptide substrates for **memapsin 2**. The substrate analogs contain at least one analog of an amide bond which is not capable of being cleaved by **memapsin 2**. Processes for the synthesis of two substrate analogues including isosteres at the sites of the critical amino acid residues were developed and the substrate analogues, OMR99-1 and OM99-2, were synthesized. OM99-2 is based on an octapeptide Glu-Val-Asn-Leu-Ala-Ala-Glu-Phe (SEQ ID NO:28) with the Leu-Ala peptide bond substituted by a transition-state **isostere** hydroxyethylene group (FIG. 1). The **inhibition** constant of OM99-2 is 1.6.times.10.sup.-9 M against **recombinant** **memapsin 2**. Crystallography of **memapsin 2** bound to this **inhibitor** was used to determine the three dimensional structure of the protein, as well as the importance of the various residues in binding. This information can be used by those skilled in the art to design new **inhibitors**, using commercially available **software programs** and techniques familiar to those in organic chemistry and enzymology, to design new **inhibitors** to **memapsin 2**, useful in **diagnostics** and for the **treatment** and/or **prevention** of **Alzheimer's** disease.

L11 ANSWER 6 OF 7 USPATFULL on STN  
AN 2002:214213 USPATFULL  
TI **Inhibitors of memapsin 2 and use thereof**  
IN Koelsch, Gerald, Oklahoma City, OK, UNITED STATES  
Tang, Jordan J.N., Edmond, OK, UNITED STATES  
Hong, Lin, Oklahoma City, OK, UNITED STATES  
Ghosh, Arun K., River Forest, IL, UNITED STATES  
PA Oklahoma Medical Research Foundation (U.S. corporation)  
PI US 2002115600 A1 20020822  
AI US 2001-845226 A1 20010430 (9)  
RLI Division of Ser. No. US 2000-603713, filed on 27 Jun 2000, PENDING  
PRAI US 1999-141363P 19990628 (60)  
US 1999-168060P 19991130 (60)  
US 2000-177836P 20000125 (60)  
US 2000-178368P 20000127 (60)  
US 2000-210292P 20000608 (60)  
DT Utility  
FS APPLICATION  
LREP Patrea L. Pabst, Arnall Golden & Gregory, LLP, 2800 One Atlantic Center,  
1201 West Peachtree Street, Atlanta, GA, 30309-3450  
CLMN Number of Claims: 23  
ECL Exemplary Claim: 1  
DRWN 12 Drawing Page(s)  
LN.CNT 2377  
CAS INDEXING IS AVAILABLE FOR THIS PATENT.  
AB Methods for the production of purified, catalytically active, recombinant **memapsin 2** have been developed.

The substrate and subsite specificity of the catalytically active enzyme have been determined. The substrate and subsite specificity information was used to design substrate analogs of the natural **memapsin** 2 substrate that can **inhibit** the function of **memapsin** 2. The substrate analogs are based on peptide sequences, shown to be related to the natural peptide substrates for **memapsin** 2. The substrate analogs contain at least one analog of an amide bond which is not capable of being cleaved by **memapsin** 2. Processes for the synthesis of two substrate analogues including isosteres at the sites of the critical amino acid residues were developed and the substrate analogues, OMR99-1 and OM99-2, were synthesized. OM99-2 is based on an octapeptide Glu-Val-Asn-Leu-Ala-Ala-Glu-Phe (SEQ ID NO:28) with the Leu-Ala peptide bond substituted by a transition-state **isostere** hydroxyethylene group (FIG. 1). The **inhibition** constant of OM99-2 is 1.6.times.10.sup.-9 M against **recombinant** **memapsin** 2. Crystallography of **memapsin** 2 bound to this **inhibitor** was used to determine the three dimensional structure of the protein, as well as the importance of the various residues in binding. This information can be used by those skilled in the art to design new **inhibitors**, using commercially available **software programs** and techniques familiar to those in organic chemistry and enzymology, to design new **inhibitors** to **memapsin** 2, useful in **diagnostics** and for the **treatment** and/or **prevention** of **Alzheimer's** disease.

L11 ANSWER 7 OF 7 USPATFULL on STN  
AN 2002:92777 USPATFULL  
TI Catalytically active **recombinant** **memapsin** and methods of use thereof  
IN Tang, Jordan J. N., Edmond, OK, UNITED STATES  
Lin, Xinli, Edmond, OK, UNITED STATES  
Koelsch, Gerald, Oklahoma City, OK, UNITED STATES  
Hong, Lin, Oklahoma City, OK, UNITED STATES  
PI US 2002049303 A1 20020425  
AI US 2001-796264 A1 20010228 (9)  
RLI Division of Ser. No. US 2000-604608, filed on 27 Jun 2000, PENDING  
PRAI US 1999-141363P 19990628 (60)  
US 1999-168060P 19991130 (60)  
US 2000-177836P 20000125 (60)  
US 2000-178368P 20000127 (60)  
DT Utility  
FS APPLICATION  
LREP PATREA L. PABST, HOLLAND & KNIGHT LLP, SUITE 2000, ONE ATLANTIC CENTER,  
1201 WEST PEACHTREE STREET, N.E., ATLANTA, GA, 30309-3400  
CLMN Number of Claims: 33  
ECL Exemplary Claim: 1  
DRWN 12 Drawing Page(s)  
LN.CNT 2441  
CAS INDEXING IS AVAILABLE FOR THIS PATENT.  
AB Methods for the production of purified, catalytically active, **recombinant** **memapsin** 2 have been developed. The substrate and subsite specificity of the catalytically active enzyme have been determined. The substrate and subsite specificity information was used to design substrate analogs of the natural **memapsin** 2 substrate that can **inhibit** the function of **memapsin** 2. The substrate analogs are based on peptide sequences, shown to be related to the natural peptide substrates for **memapsin** 2. The substrate analogs contain at least one analog of an amide bond which is not capable of being cleaved by **memapsin** 2. Processes for the synthesis of two substrate analogs including isosteres at the sites of the critical amino acid residues were developed and the substrate analogs, OMR99-1 and OM99-2, were synthesized. OM99-2 is based on an octapeptide

Glu-Val-Asn-Leu-Ala-Ala-Glu-Phe (SEQ ID NO:28) with the Leu-Ala peptide bond substituted by a transition-state **isostere** hydroxyethylene group (FIG. 1). The **inhibition** constant of OM99-2 is 1.6.times.10.<sup>-9</sup> M against **recombinant** pro-**memapsin** 2. Crystallography of **memapsin** 2 bound to this **inhibitor** was used to determine the three dimensional structure of the protein, as well as the importance of the various residues in binding. This information can be used by those skilled in the art to design new **inhibitors**, using commercially available **software programs** and techniques familiar to those in organic chemistry and enzymology, to design new **inhibitors** to **memapsin** 2, useful in **diagnostics** and for the **treatment** and/or **prevention** of **Alzheimer's** disease.

=> d his

(FILE 'HOME' ENTERED AT 16:27:26 ON 05 AUG 2003)

FILE 'MEDLINE, CAPLUS, BIOSIS, BIOTECHDS, EMBASE, USPATFULL, WPIDS'  
ENTERED AT 16:29:30 ON 05 AUG 2003

L1 630677 S (MEMAPSIN 2 OR BETA SECRETASE OR BETA AMYLOID PRECURSOR PROTE  
L2 135 S (L1 OR MEMAPSIN 2) AND (OM 99 OR OM-99 OR DIPEPTIDE ISOSTERE  
L3 22 S L2 AND (COMPUTER PROGRAM# OR SOFTWARE PROGRAM#)  
L4 22 S L3 AND (RECOMBINANT?)  
L5 22 S L4 AND (TREAT? OR THERAPEUT? OR DIAGNOS? OR PREVENT?)  
L6 22 S L5 AND (INHIBIT?)  
L7 8203 S TANG J?/AU  
L8 417 S HONG L/AU  
L9 1933 S GHOSH A/AU  
L10 100 S KOELSCH G/AU  
L11 7 S L6 AND (L7 OR L8 OR L9 OR L10)

=> d 16 1-22 bib ab

L6 ANSWER 1 OF 22 CAPLUS COPYRIGHT 2003 ACS on STN  
AN 2001:12489 CAPLUS  
DN 134:80832  
TI **Inhibitors of memapsin 2 and use thereof**  
IN Tang, Jordan J. N.; Hong, Ling; Ghosh, Arun K.  
PA Oklahoma Medical Research Foundation, USA; The Board of Trustees of the University of Illinois  
SO PCT Int. Appl., 86 pp.  
CODEN: PIXXD2  
DT Patent  
LA English

FAN.CNT 3

	PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
PI	WO 2001000665	A2	20010104	WO 2000-US17742	20000627
	WO 2001000665	A3	20010927		
	WO 2001000665	C2	20020725		
		W:	AE, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CR, CU, CZ, DE, DK, DM, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, UZ, VN, YU, ZA, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM		
		RW:	GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG		
EP	1194449	A2	20020410	EP 2000-943236	20000627
	R:	AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT, IE, SI, LT, LV, FI, RO			

JP 2003506322	T2	20030218	JP 2001-507071	20000627
US 6545127	B1	20030408	US 2000-604608	20000627
US 2002049303	A1	20020425	US 2001-796264	20010228
US 2002164760	A1	20021107	US 2001-795903	20010228
US 2002115600	A1	20020822	US 2001-845226	20010430
PRAI US 1999-141363P	P	19990628		
US 1999-168060P	P	19991130		
US 2000-177836P	P	20000125		
US 2000-178368P	P	20000127		
US 2000-210292P	P	20000608		
US 2000-603713	A3	20000627		
US 2000-604608	A3	20000627		
WO 2000-US17742	W	20000627		

AB Methods for the prodn. of purified, catalytically active, recombinant memapsin 2 have been developed. The substrate and subsite specificity of the catalytically active enzyme have been detd. The substrate and subsite specificity information was used to design substrate analogs of the natural memapsin 2 substrate that can inhibit the function of memapsin 2. The substrate analogs are based on peptide sequences, shown to be related to the natural peptide substrates for memapsin 2. The substrate analogs contain at least one analog of an amide bond which is not capable of being cleaved by memapsin 2. Processes for the synthesis of two substrate analogs including isosteres at the sites of the crit. amino acid residues were developed and the substrate analogs, OMR99-1 and OM99-2, were synthesized. OM99-2 is based on an octapeptide Glu-Val-Asn-Leu-Ala-Ala-Glu-Phe (SEQ ID NO:28) with the Leu-Ala peptide bond substituted by a transition-state isostere hydroxyethylene group (Figure 1). The inhibition const. of OM99-2 is 1.6 x 10-9 M against recombinant pro-memapsin 2. Crystallog. of memapsin 2 bound to this inhibitor was used to det. the three dimensional structure of the protein, as well as the importance of the various residues in binding. This information can be used by those skilled in the art to design new inhibitors, using com. available software programs and techniques familiar to those in org. chem. and enzymol., to design new inhibitors to memapsin 2, useful in diagnostics and for the treatment and/or prevention of Alzheimer's disease.

L6 ANSWER 2 OF 22 CAPLUS COPYRIGHT 2003 ACS on STN

AN 2001:12487 CAPLUS

DN 134:68049

TI Catalytically active recombinant memapsin 2, 3D crystal structure based inhibitor design, synthesis, and screening, for Alzheimer's disease treatment

IN Tang, Jordan J. N.; Lin, Xinli; Koelsch, Gerald

PA Oklahoma Medical Research Foundation, USA

SO PCT Int. Appl., 87 pp.

CODEN: PIXXD2

DT Patent

LA English

FAN.CNT 3

	PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
	-----	-----	-----	-----	-----
PI	WO 2001000663	A2	20010104	WO 2000-US17661	20000627
	WO 2001000663	A3	20011004		
	W: AE, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CR, CU, CZ, DE, DK, DM, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, UZ, VN, YU, ZA, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM				
	RW: GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY,				

DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF, BJ,  
CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG  
EP 1196609 A2 20020417 EP 2000-943208 20000627  
R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT,  
IE, SI, LT, LV, FI, RO  
JP 2003503072 T2 20030128 JP 2001-507069 20000627  
US 6545127 B1 20030408 US 2000-604608 20000627  
US 2002049303 A1 20020425 US 2001-796264 20010228  
US 2002164760 A1 20021107 US 2001-795903 20010228  
US 2002115600 A1 20020822 US 2001-845226 20010430  
PRAI US 1999-141363P P 19990628  
US 1999-168060P P 19991130  
US 2000-177836P P 20000125  
US 2000-178368P P 20000127  
US 2000-210292P P 20000608  
US 2000-603713 A3 20000627  
US 2000-604608 A3 20000627  
WO 2000-US17661 W 20000627  
AB A method for producing catalytically active **recombinant memapsin 2** comprising expression in a bacteria and refolding the **recombinant memapsin 2** under conditions which dissociate and then slowly refold the enzyme into a catalytically active form is disclosed. A method of isolating **inhibitors of cleavage by memapsin 2** comprising adding to one or more potential **inhibitors** of catalytically active **recombinant memapsin 2**, and a substrate for **memapsin 2**, and screening for decreased cleavage of the substrate by the **inhibitors**, wherein the **inhibitors** are in a library of small synthetic mols., like proteins and peptides. Alternatively, the **inhibitors** are oligonucleotides preventing or decreasing expression of catalytically active **memapsin 2**. A method for designing or obtaining **inhibitors** of catalytically active **memapsin 2** comprising modeling an **inhibitor** based on the crystn. coordinates of **memapsin 2** or parameters. A database comprising binding properties and chem. structures of compds. designed or screened by modeling an **inhibitor** based on the crystn. coordinates of **memapsin 2** or parameters is claimed. A method of **treating or preventing** Alzheimer's disease comprising administering to a patient in need thereof an **inhibitor of memapsin 2** which binds to the active site of the **memapsin 2** defined by the presence of two catalytic aspartic residues and substrate binding cleft, is also claimed. The cDNAs of two new human membrane-assocd. aspartic proteases, **memapsin 1** and **memapsin 2**, have been cloned and sequenced. The substrate and subsite specificity of the catalytically active enzyme have been detd. The substrate and subsite specificity information was used to design substrate analogs of the natural **memapsin 2** substrate that can **inhibit** the function of **memapsin 2**. The substrate analogs are based on peptide sequences, shown to be related to the natural peptide substrates for **memapsin 2**. The substrate analogs contain at least one analog of an amide bond which is not capable of being cleaved by **memapsin 2**. Processes for the synthesis of two substrate analogs including isosteres at the sites of the crit. amino acid residues were developed and the substrate analogs, OMR99-1 and OM99-2, were synthesized. OM99-2 is based on an octapeptide Glu-Val-Asn-Leu-Ala-Ala-Glu-Phe (SEQ ID NO:28) with the Leu-Ala peptide bond substituted by a transition-state **isostere** hydroxyethylene group (Fig. 1). The **inhibition const.** of OM99-2 is 1.6 x 10<sup>9</sup> M against **recombinant pro-memapsin 2**. Crystallog. of **memapsin 2** bound to this **inhibitor** was used to det. the three dimensional structure of the protein, as well as the importance of the various residues in binding. This information can be used to design new **inhibitors**, using

com. available **software programs** and techniques familiar to those in org. chem. and enzymol., to design new **inhibitors to memapsin 2**, useful in **diagnostics** and for the **treatment** and/or **prevention** of **Alzheimer's disease**.

L6 ANSWER 3 OF 22 USPATFULL on STN  
AN 2003:200784 USPATFULL  
TI Immunogenic HBC chimer particles having enhanced stability  
IN Birkett, Ashley J., Escondido, CA, UNITED STATES  
PI US 2003138769 A1 20030724  
AI US 2001-930915 A1 20010815 (9)  
RLI Continuation-in-part of Ser. No. US 2000-226867, filed on 22 Aug 2000, PENDING Continuation-in-part of Ser. No. US 2000-225843, filed on 16 Aug 2000, PENDING  
DT Utility  
FS APPLICATION  
LREP WELSH & KATZ, LTD, 120 S RIVERSIDE PLAZA, 22ND FLOOR, CHICAGO, IL, 60606  
CLMN Number of Claims: 115  
ECL Exemplary Claim: 1  
DRWN 10 Drawing Page(s)  
LN.CNT 6993  
AB A chimeric, carboxy-terminal truncated hepatitis B virus nucleocapsid protein (HBC) is disclosed that is engineered for both enhanced stability of self-assembled particles and the display of an immunogenic epitope. The display of the immunogenic epitope is displayed in the immunogenic loop of HBC, whereas the enhanced stability of self-assembled particles is obtained by the presence of at least one heterologous cysteine residue near the carboxy-terminus of the chimer molecule. Methods of making and using the chimers are also disclosed.

L6 ANSWER 4 OF 22 USPATFULL on STN  
AN 2003:187895 USPATFULL  
TI 12 human secreted proteins  
IN Ni, Jian, Germantown, MD, UNITED STATES  
Young, Paul E., Gaithersburg, MD, UNITED STATES  
Kenny, Joseph J., Damascus, MD, UNITED STATES  
Olsen, Henrik S., Gaithersburg, MD, UNITED STATES  
Moore, Paul A., Germantown, MD, UNITED STATES  
Wei, Ying-Fei, Berkeley, CA, UNITED STATES  
Greene, John M., Gaithersburg, MD, UNITED STATES  
Ruben, Steven M., Olney, MD, UNITED STATES  
PI US 2003129685 A1 20030710  
AI US 2001-836353 A1 20010418 (9)  
RLI Continuation-in-part of Ser. No. WO 1999-US25031, filed on 27 Oct 1999, UNKNOWN  
PRAI US 1998-105971P 19981028 (60)  
US 2000-198407P 20000419 (60)

DT Utility  
FS APPLICATION  
LREP HUMAN GENOME SCIENCES INC, 9410 KEY WEST AVENUE, ROCKVILLE, MD, 20850  
CLMN Number of Claims: 23  
ECL Exemplary Claim: 1  
DRWN 59 Drawing Page(s)  
LN.CNT 31945

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB The present invention relates to novel human secreted proteins and isolated nucleic acids containing the coding regions of the genes encoding such proteins. Also provided are vectors, host cells, antibodies, and recombinant methods for producing human secreted proteins. The invention further relates to diagnostic and therapeutic methods useful for diagnosing and treating diseases, disorders, and/or conditions related to these novel human secreted proteins.

L6 ANSWER 5 OF 22 USPATFULL on STN  
AN 2003:165862 USPATFULL  
TI Directed evolution of novel binding proteins  
IN Ladner, Robert Charles, Ijamsville, MD, UNITED STATES  
Guterman, Sonia Kosow, Belmont, MA, UNITED STATES  
Roberts, Bruce Lindsay, Milford, MA, UNITED STATES  
Markland, William, Milford, MA, UNITED STATES  
Ley, Arthur Charles, Newton, MA, UNITED STATES  
Kent, Rachel Baribault, Boxborough, MA, UNITED STATES  
PI US 2003113717 A1 20030619  
AI US 2001-893878 A1 20010629 (9)  
RLI Continuation of Ser. No. US 1997-993776, filed on 18 Dec 1997, PENDING  
Continuation of Ser. No. US 1995-415922, filed on 3 Apr 1995, PATENTED  
Continuation of Ser. No. US 1993-9319, filed on 26 Jan 1993, PATENTED  
Division of Ser. No. US 1991-664989, filed on 1 Mar 1991, PATENTED  
Continuation-in-part of Ser. No. US 1990-487063, filed on 2 Mar 1990,  
ABANDONED Continuation-in-part of Ser. No. US 1988-240160, filed on 2  
Sep 1988, ABANDONED  
PRAI WO 1989-US3731 19890901  
DT Utility  
FS APPLICATION  
LREP BROWDY AND NEIMARK, P.L.L.C., 624 Ninth Street, N.W., Washington, DC,  
20001  
CLMN Number of Claims: 25  
ECL Exemplary Claim: 1  
DRWN 16 Drawing Page(s)  
LN.CNT 15933  
CAS INDEXING IS AVAILABLE FOR THIS PATENT.  
AB In order to obtain a novel binding protein against a chosen target, DNA  
molecules, each encoding a protein comprising one of a family of similar  
potential binding domains and a structural-signal calling for the  
display of the protein on the outer surface of a chosen bacterial cell,  
bacterial spore or phage (genetic package) are introduced into a genetic  
package. The protein is expressed and the potential binding domain is  
displayed on the outer surface of the package. The cells or viruses  
bearing the binding domains which recognize the target molecule are  
isolated and amplified. The successful binding domains are then  
characterized. One or more of these successful binding domains is used  
as a model for the design of a new family of potential binding domains,  
and the process is repeated until a novel binding domain having a  
desired affinity for the target molecule is obtained. In one embodiment,  
the first family of potential binding domains is related to bovine  
pancreatic trypsin inhibitor, the genetic package is M13  
phage, and the protein includes the outer surface transport signal of  
the M13 gene III protein.

L6 ANSWER 6 OF 22 USPATFULL on STN  
AN 2003:134541 USPATFULL  
TI Inhibitors of memapsin 2 and use thereof  
IN Tang, Jordan J. N., Edmond, OK, UNITED STATES  
Koelsch, Gerald, Oklahoma City, OK, UNITED STATES  
Ghosh, Arun K., River Forest, IL, UNITED STATES  
PA Oklahoma Medical Research Foundation, Oklahoma City, OK (U.S.  
corporation)  
PI US 2003092629 A1 20030515  
AI US 2001-32818 A1 20011228 (10)  
PRAI US 2001-275756P 20010314 (60)  
US 2000-258705P 20001228 (60)  
DT Utility  
FS APPLICATION  
LREP HAMILTON, BROOK, SMITH & REYNOLDS, P.C., 530 VIRGINIA ROAD, P.O: BOX  
9133, CONCORD, MA, 01742-9133  
CLMN Number of Claims: 24  
ECL Exemplary Claim: 1  
DRWN 9 Drawing Page(s)

LN.CNT 2203

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB Methods for the production of purified, catalytically active, recombinant memapsin 2 have been developed-. The substrate and subsite specificity of the catalytically active enzyme have been determined by a method which determines the initial hydrolysis rate of the substrates by using MALDI-TOF/MS. Alternatively, the subsite specificity of memapsin can be determined by probing a library of inhibitors with memapsin 2 and subsequently detecting the bound memapsin 2 with an antibody raised to memapsin 2 and an alkaline phosphatase conjugated secondary antibody. The substrate and subsite specificity information was used to design substrate analogs of the natural memapsin 2 substrate that can inhibit the function of memapsin 2. The substrate analogs are based on peptide sequences, shown to be related to the natural peptide substrates for memapsin 2. The substrate analogs contain at least one analog of an amide bond which is not capable of being cleaved by memapsin 2. Processes for the synthesis of substrate analogues including isosteres at the sites of the critical amino acid residues were developed and the more than seventy substrate analogues were synthesized, among which MMI-005, MMI-012, MMI-017, MMI-018, MMI-025, MMI-026, MMI-037, MMI-039, MMI-040, MMI-066, MMI-070, and MMI-071 have inhibition constants in the range of 1.4-61.4.times.10.sup.-9 M against recombinant pro-memapsin 2. These inhibitors are useful in diagnostics and for the treatment and/or prevention of Alzheimer's disease.

L6 ANSWER 7 OF 22 USPATFULL on STN

AN 2003:96167 USPATFULL

TI Catalytically active recombinant memapsin and methods of use thereof

IN Tang, Jordan J. N., Edmond, OK, United States

Lin, Xinli, Edmond, OK, United States

Koelsch, Gerald, Oklahoma City, OK, United States

Hong, Lin, Oklahoma City, OK, United States

PA Oklahoma Medical Research Foundation, Oklahoma City, OK, United States (U.S. corporation)

PI US 6545127 B1 20030408

AI US 2000-604608 20000627 (9)

PRAI US 1999-141363P 19990628 (60)

US 1999-168060P 19991130 (60)

US 2000-177836P 20000125 (60)

US 2000-178368P 20000127 (60)

US 2000-210292P 20000608 (60)

DT Utility

FS GRANTED

EXNAM Primary Examiner: Borin, Michael; Assistant Examiner: Zhou, Shuba

LREP Hamilton, Brook, Smith & Reynolds, P.C.

CLMN Number of Claims: 18

ECL Exemplary Claim: 7

DRWN 21 Drawing Figure(s); 12 Drawing Page(s)

LN.CNT 2563

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB Methods for the production of purified, catalytically active, recombinant memapsin 2 have been developed.

The substrate and subsite specificity of the catalytically active enzyme have been determined. The substrate and subsite specificity information was used to design substrate analogs of the natural memapsin 2 substrate that can inhibit the function of

memapsin 2. The substrate analogs are based on peptide sequences, shown to be related to the natural peptide substrates for memapsin 2. The substrate analogs contain at least one analog of an amide bond which is not capable of being cleaved by

**memapsin 2.** Processes for the synthesis of two substrate analogues including isosteres at the sites of the critical amino acid residues were developed and the substrate analogues, OMR99-1 and OM99-2, were synthesized. OM99-2 is based on an octapeptide Glu-Val-Asn-Leu-Ala-Ala-Glu-Phe (SEQ ID NO:28) with the Leu-Ala peptide bond substituted by a transition-state **isostere** hydroxyethylene group (FIG. 1). The **inhibition** constant of OM99-2 is 1.6.times.10.<sup>sup.-9M</sup> against **recombinant** **memapsin 2.** **Crystallography** of **memapsin 2** bound to this **inhibitor** was used to determine the three dimensional structure of the protein, as well as the importance of the various residues in binding. This information can be used by those skilled in the art to design new **inhibitors**, using commercially available **software programs** and techniques familiar to those in organic chemistry and enzymology, to design new **inhibitors** to **memapsin 2**, useful in **diagnostics** and for the **treatment** and/or **prevention** of **Alzheimer's** disease.

L6 ANSWER 8 OF 22 USPATFULL on STN  
AN 2003:79303 USPATFULL  
TI 12 human secreted proteins  
IN Ni, Jian, Germantown, MD, UNITED STATES  
Young, Paul E., Gaithersburg, MD, UNITED STATES  
Kenny, Joseph J., Damascus, MD, UNITED STATES  
Olsen, Henrik S., Gaithersburg, MD, UNITED STATES  
Moore, Paul A., Germantown, MD, UNITED STATES  
Wei, Ying-Fei, Berkeley, CA, UNITED STATES  
Greene, John M., Gaithersburg, MD, UNITED STATES  
Ruben, Steven M., Olney, MD, UNITED STATES  
Liu, Ding, Gaithersburg, MD, UNITED STATES  
Crocker, Paul R., Dundee, UNITED KINGDOM  
PI US 2003055231 A1 20030320  
AI US 2001-984130 A1 20011029 (9)  
RLI Continuation-in-part of Ser. No. US 2001-836353, filed on 18 Apr 2001,  
PENDING Continuation-in-part of Ser. No. WO 1999-US25031, filed on 27  
Oct 1999, UNKNOWN  
PRAI US 2000-243792P 20001030 (60)  
US 2000-198407P 20000419 (60)  
US 1998-105971P 19981028 (60)  
DT Utility  
FS APPLICATION  
LREP HUMAN GENOME SCIENCES INC, 9410 KEY WEST AVENUE, ROCKVILLE, MD, 20850  
CLMN Number of Claims: 23  
ECL Exemplary Claim: 1  
DRWN 67 Drawing Page(s)  
LN.CNT 31982  
CAS INDEXING IS AVAILABLE FOR THIS PATENT.  
AB The present invention relates to 12 novel human secreted proteins and isolated nucleic acids containing the coding regions of the genes encoding such proteins. Also provided are vectors, host cells, antibodies, and **recombinant** methods for producing human secreted proteins. The invention further relates to **diagnostic** and **therapeutic** methods useful for **diagnosing** and **treating** disorders related to these novel human secreted proteins.

L6 ANSWER 9 OF 22 USPATFULL on STN  
AN 2003:4068 USPATFULL  
TI Method of **preventing** cell death using segments of neural thread proteins  
IN Averback, Paul A., Beaconsfield, CANADA  
PI US 2003004107 A1 20030102  
AI US 2002-146130 A1 20020516 (10)  
PRAI US 2001-290971P 20010516 (60)

DT Utility  
FS APPLICATION  
LREP HUNTON & WILLIAMS, INTELLECTUAL PROPERTY DEPARTMENT, 1900 K STREET,  
N.W., SUITE 1200, WASHINGTON, DC, 20006-1109  
CLMN Number of Claims: 14  
ECL Exemplary Claim: 1  
DRWN 9 Drawing Page(s)  
LN.CNT 1698  
CAS INDEXING IS AVAILABLE FOR THIS PATENT.  
AB Disclosed is a method of preventing, inhibiting,  
and/or ameliorating cell death and/or tissue necrosis in live tissue by  
contacting live tissue with at least a segment of NTP, or homologue,  
variant, derivative or mimetic thereof, where the segment of NTP, or  
homologue, variant, derivative or mimetic thereof is present in an  
amount effective to prevent, inhibit, and/or  
ameliorate cell death and/or tissue necrosis. The method is capable of  
treating conditions requiring prevention,  
inhibition, and/or amelioration of cell death and/or tissue  
necrosis.

L6 ANSWER 10 OF 22 USPATFULL on STN  
AN 2003:3410 USPATFULL  
TI Method of preventing cell death using antibodies to neural  
thread proteins  
IN Averback, Paul A., Quebec, CANADA  
PI US 2003003445 A1 20030102  
AI US 2002-138516 A1 20020506 (10)  
PRAI US 2001-288463P 20010504 (60)  
DT Utility  
FS APPLICATION  
LREP HUNTON & WILLIAMS, INTELLECTUAL PROPERTY DEPARTMENT, 1900 K STREET,  
N.W., SUITE 1200, WASHINGTON, DC, 20006-1109  
CLMN Number of Claims: 12  
ECL Exemplary Claim: 1  
DRWN 10 Drawing Page(s)  
LN.CNT 1705  
CAS INDEXING IS AVAILABLE FOR THIS PATENT.  
AB Disclosed is a method of preventing, inhibiting,  
and/or ameliorating cell death and/or tissue necrosis in live tissue  
containing neural thread proteins (NTP) by contacting the live tissue  
with at least an antibody, antibody fragment or antibody derivative that  
recognizes or binds to NTP, where the antibody, antibody fragment or  
antibody derivative is present in an amount effective to prevent  
, inhibit, reduce, control and/or ameliorate cell death and/or  
tissue necrosis. The method is capable of treating conditions  
requiring prevention, inhibition, reduction, control  
and/or amelioration of cell death and/or tissue necrosis caused by the  
presence of NTP.

L6 ANSWER 11 OF 22 USPATFULL on STN  
AN 2002:294717 USPATFULL  
TI Catalytically active recombinant memapsin and methods of use  
thereof  
IN Lin, Xinli, Edmond, OK, UNITED STATES  
Koelsch, Gerald, Oklahoma City, OK, UNITED STATES  
Tang, Jordan J.N., Edmond, OK, UNITED STATES  
PA Oklahoma Medical Research Foundation  
PI US 2002164760 A1 20021107  
AI US 2001-795903 A1 20010228 (9)  
RLI Division of Ser. No. US 2000-604608, filed on 27 Jun 2000, PENDING  
PRAI US 1999-141363P 19990628 (60)  
US 1999-168060P 19991130 (60)  
US 2000-177836P 20000125 (60)  
US 2000-178368P 20000127 (60)  
US 2000-210292P 20000608 (60)

DT Utility  
FS APPLICATION  
LREP PATREA L. PABST, HOLLAND & KNIGHT LLP, SUITE 2000, ONE ATLANTIC CENTER,  
1201 WEST PEACHTREE STREET, N.E., ATLANTA, GA, 30309-3400  
CLMN Number of Claims: 33  
ECL Exemplary Claim: 1  
DRWN 12 Drawing Page(s)  
LN.CNT 2440

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB Methods for the production of purified, catalytically active, recombinant memapsin 2 have been developed. The substrate and subsite specificity of the catalytically active enzyme have been determined. The substrate and subsite specificity information was used to design substrate analogs of the natural memapsin 2 substrate that can inhibit the function of memapsin 2. The substrate analogs are based on peptide sequences, shown to be related to the natural peptide substrates for memapsin 2. The substrate analogs contain at least one analog of an amide bond which is not capable of being cleaved by memapsin 2. Processes for the synthesis of two substrate analogues including isosteres at the sites of the critical amino acid residues were developed and the substrate analogues, OMR99-1 and OM99-2, were synthesized. OM99-2 is based on an octapeptide Glu-Val-Asn-Leu-Ala-Ala-Glu-Phe (SEQ ID NO:28) with the Leu-Ala peptide bond substituted by a transition-state isostere hydroxyethylene group (FIG. 1). The inhibition constant of OM99-2 is 1.6.times.10.sup.-9 M against recombinant pro-memapsin 2. Crystallography of memapsin 2 bound to this inhibitor was used to determine the three dimensional structure of the protein, as well as the importance of the various residues in binding. This information can be used by those skilled in the art to design new inhibitors, using commercially available software programs and techniques familiar to those in organic chemistry and enzymology, to design new inhibitors to memapsin 2, useful in diagnostics and for the treatment and/or prevention of Alzheimer's disease.

L6 ANSWER 12 OF 22 USPATFULL on STN  
AN 2002:272761 USPATFULL  
TI Directed evolution of novel binding proteins  
IN Ladner, Robert Charles, Ijamsville, MD, UNITED STATES  
Guterman, Sonia Kosow, Belmont, MA, UNITED STATES  
Roberts, Bruce Lindsay, Milford, MA, UNITED STATES  
Markland, William, Milford, MA, UNITED STATES  
Ley, Arthur Charles, Newton, MA, UNITED STATES  
Kent, Rachel Baribault, Boxborough, MA, UNITED STATES  
PI US 2002150881 A1 20021017  
AI US 2001-781988 A1 20010214 (9)  
RLI Continuation of Ser. No. US 1998-192067, filed on 16 Nov 1998, ABANDONED  
Continuation of Ser. No. US 1995-415922, filed on 3 Apr 1995, PATENTED  
Continuation of Ser. No. US 1993-9319, filed on 26 Jan 1993, PATENTED  
Division of Ser. No. US 1991-664989, filed on 1 Mar 1991, PATENTED  
Continuation-in-part of Ser. No. US 1990-487063, filed on 2 Mar 1990,  
ABANDONED Continuation-in-part of Ser. No. US 1988-240160, filed on 2  
Sep 1988, ABANDONED  
PRAI WO 1989-US3731 19890901  
DT Utility  
FS APPLICATION  
LREP BROWDY AND NEIMARK, P.L.L.C., 624 Ninth Street, N.W., Washington, DC,  
20001  
CLMN Number of Claims: 18  
ECL Exemplary Claim: 1  
DRWN 16 Drawing Page(s)  
LN.CNT 15696

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB In order to obtain a novel binding protein against a chosen target, DNA molecules, each encoding a protein comprising one of a family of similar potential binding domains and a structural signal calling for the display of the protein on the outer surface of a chosen bacterial cell, bacterial spore or phage (genetic package) are introduced into a genetic package. The protein is expressed and the potential binding domain is displayed on the outer surface of the package. The cells or viruses bearing the binding domains which recognize the target molecule are isolated and amplified. The successful binding domains are then characterized. One or more of these successful binding domains is used as a model for the design of a new family of potential binding domains, and the process is repeated until a novel binding domain having a desired affinity for the target molecule is obtained. In one embodiment, the first family of potential binding domains is related to bovine pancreatic trypsin inhibitor, the genetic package is M13 phage, and the protein includes the outer surface transport signal of the M13 gene III protein.

L6 ANSWER 13 OF 22 USPATFULL on STN

AN 2002:265848 USPATFULL

TI Biopolymer sequence comparison

IN Toll, Lawrence R., Redwood City, CA, UNITED STATES

Lincoln, Patrick Denis, Woodside, CA, UNITED STATES

Karp, Peter, San Mateo, CA, UNITED STATES

Sonmez, Kemal, Menlo Park, CA, UNITED STATES

PI US 2002146724 A1 20021010

AI US 2001-6492 A1 20011203 (10)

PRAI US 2000-250743P 20001201 (60)

DT Utility

FS APPLICATION

LREP DAVID L. FEIGENBAUM, Fish & Richardson P.C., 225 Franklin Street, Boston, MA, 02110-2804

CLMN Number of Claims: 71

ECL Exemplary Claim: 1

DRWN 7 Drawing Page(s)

LN.CNT 1796

AB Disclosed are methods, software, and systems for comparing biopolymer sequences. The model includes at least two different characterizations of states of matching between segments of sequences at defined positions. Examples of states of matching include: similarity and dissimilarity between objects, as well as similarity to a reference, e.g., a reference sequence or a sequence profile. A topology of particular match states can be used to identify classes of sequences, e.g., preprohormone sequences.

L6 ANSWER 14 OF 22 USPATFULL on STN

AN 2002:214213 USPATFULL

TI Inhibitors of memapsin 2 and use thereof

IN Koelsch, Gerald, Oklahoma City, OK, UNITED STATES

Tang, Jordan J.N., Edmond, OK, UNITED STATES

Hong, Lin, Oklahoma City, OK, UNITED STATES

Ghosh, Arun K., River Forest, IL, UNITED STATES

PA Oklahoma Medical Research Foundation (U.S. corporation)

PI US 2002115600 A1 20020822

AI US 2001-845226 A1 20010430 (9)

RLI Division of Ser. No. US 2000-603713, filed on 27 Jun 2000, PENDING

PRAI US 1999-141363P 19990628 (60)

US 1999-168060P 19991130 (60)

US 2000-177836P 20000125 (60)

US 2000-178368P 20000127 (60)

US 2000-210292P 20000608 (60)

DT Utility

FS APPLICATION

LREP Patrea L. Pabst, Arnall Golden & Gregory, LLP, 2800 One Atlantic Center,

1201 West Peachtree Street, Atlanta, GA, 30309-3450

CLMN Number of Claims: 23

ECL Exemplary Claim: 1

DRWN 12 Drawing Page(s)

LN.CNT 2377

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB Methods for the production of purified, catalytically active, recombinant memapsin 2 have been developed.

The substrate and subsite specificity of the catalytically active enzyme have been determined. The substrate and subsite specificity information was used to design substrate analogs of the natural memapsin 2 substrate that can inhibit the function of

memapsin 2. The substrate analogs are based on peptide sequences, shown to be related to the natural peptide substrates for

memapsin 2. The substrate analogs contain at least one analog of an amide bond which is not capable of being cleaved by

memapsin 2. Processes for the synthesis of two substrate analogues including isosteres at the sites of the critical amino acid residues were developed and the substrate analogues, OMR99-1 and OM99-2, were synthesized. OM99-2 is based on an octapeptide

Glu-Val-Asn-Leu-Ala-Ala-Glu-Phe (SEQ ID NO:28) with the Leu-Ala peptide bond substituted by a transition-state isostere

hydroxyethylene group (FIG. 1). The inhibition constant of

OM99-2 is 1.6.times.10.sup.-9 M against recombinant pro-

memapsin 2. Crystallography of memapsin

2 bound to this inhibitor was used to determine the

three dimensional structure of the protein, as well as the importance of the various residues in binding. This information can be used by

those skilled in the art to design new inhibitors, using

commercially available software programs and

techniques familiar to those in organic chemistry and enzymology, to

design new inhibitors to memapsin 2,

useful in diagnostics and for the treatment and/or

prevention of Alzheimer's disease.

L6 ANSWER 15 OF 22 USPATFULL on STN

AN 2002:191539 USPATFULL

TI Full-length human cDNAs encoding potentially secreted proteins

IN Milne Edwards, Jean-Baptiste Dumas, Paris, FRANCE

Bougueret, Lydie, Petit Lancy, SWITZERLAND

Jobert, Severin, Paris, FRANCE

PI US 2002102604 A1 20020801

AI US 2000-731872 A1 20001207 (9)

PRAI US 1999-169629P 19991208 (60)

US 2000-187470P 20000306 (60)

DT Utility

FS APPLICATION

LREP John Lucas, Ph.D., J.D., Genset Corporation, 10665 Sorrento Valley Road, San Diego, CA, 92121-1609

CLMN Number of Claims: 29

ECL Exemplary Claim: 1

DRWN 5 Drawing Page(s)

LN.CNT 28061

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB The invention concerns GENSET polynucleotides and polypeptides. Such GENSET products may be used as reagents in forensic analyses, as chromosome markers, as tissue/cell/organelle-specific markers, in the production of expression vectors. In addition, they may be used in screening and diagnosis assays for abnormal GENSET expression and/or biological activity and for screening compounds that may be used in the treatment of GENSET-related disorders.

L6 ANSWER 16 OF 22 USPATFULL on STN

AN 2002:92777 USPATFULL

TI Catalytically active recombinant memapsin and methods of use

thereof

IN Tang, Jordan J. N., Edmond, OK, UNITED STATES  
Lin, Xinli, Edmond, OK, UNITED STATES  
Koelsch, Gerald, Oklahoma City, OK, UNITED STATES  
Hong, Lin, Oklahoma City, OK, UNITED STATES

PI US 2002049303 A1 20020425

AI US 2001-796264 A1 20010228 (9)

RLI Division of Ser. No. US 2000-604608, filed on 27 Jun 2000, PENDING

PRAI US 1999-141363P 19990628 (60)  
US 1999-168060P 19991130 (60)  
US 2000-177836P 20000125 (60)  
US 2000-178368P 20000127 (60)

DT Utility

FS APPLICATION

LREP PATREA L. PABST, HOLLAND & KNIGHT LLP, SUITE 2000, ONE ATLANTIC CENTER,  
1201 WEST PEACHTREE STREET, N.E., ATLANTA, GA, 30309-3400

CLMN Number of Claims: 33

ECL Exemplary Claim: 1

DRWN 12 Drawing Page(s)

LN.CNT 2441

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB Methods for the production of purified, catalytically active, recombinant memapsin 2 have been developed. The substrate and subsite specificity of the catalytically active enzyme have been determined. The substrate and subsite specificity information was used to design substrate analogs of the natural memapsin 2 substrate that can inhibit the function of memapsin 2. The substrate analogs are based on peptide sequences, shown to be related to the natural peptide substrates for memapsin 2. The substrate analogs contain at least one analog of an amide bond which is not capable of being cleaved by memapsin 2. Processes for the synthesis of two substrate analogs including isosteres at the sites of the critical amino acid residues were developed and the substrate analogs, OMR99-1 and OM99-2, were synthesized. OM99-2 is based on an octapeptide Glu-Val-Asn-Leu-Ala-Ala-Glu-Phe (SEQ ID NO:28) with the Leu-Ala peptide bond substituted by a transition-state isostere hydroxyethylene group (FIG. 1). The inhibition constant of OM99-2 is 1.6.times.10.sup.-9 M against recombinant pro-memapsin 2. Crystallography of memapsin 2 bound to this inhibitor was used to determine the three dimensional structure of the protein, as well as the importance of the various residues in binding. This information can be used by those skilled in the art to design new inhibitors, using commercially available software programs and techniques familiar to those in organic chemistry and enzymology, to design new inhibitors to memapsin 2, useful in diagnostics and for the treatment and/or prevention of Alzheimer's disease.

L6 ANSWER 17 OF 22 USPATFULL on STN  
AN 2000:7289 USPATFULL  
TI Human nucleic acid binding protein  
IN Bandman, Olga, Mountain View, CA, United States  
Au-Young, Janice, Berkeley, CA, United States  
Hawkins, Phillip R., Mountain View, CA, United States  
Hillman, Jennifer L., San Jose, CA, United States  
PA Incyte Pharmaceuticals, Inc., Palo Alto, CA, United States (U.S. corporation)  
PI US 6015788 20000118  
AI US 1998-195855 19981119 (9)  
RLI Division of Ser. No. US 1996-698407, filed on 15 Aug 1996, now patented, Pat. No. US 5856128  
DT Utility  
FS Granted

EXNAM Primary Examiner: Sisson, Bradley; Assistant Examiner: Longton, Enrique D.  
LREP Sather, Susan K. Incyte Pharmaceuticals, Inc.  
CLMN Number of Claims: 2  
ECL Exemplary Claim: 1  
DRWN 11 Drawing Figure(s); 11 Drawing Page(s)  
LN.CNT 1830  
CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB The present invention provides polynucleotides which identify and encode a novel human nucleic acid binding protein (NABP). The invention provides for genetically engineered expression vectors and host cells comprising the nucleic acid sequences encoding NABP. The invention also provides for the use of substantially purified NABP or its antagonists, in pharmaceutical compositions for the treatment of diseases associated with the expression of NABP. Additionally, the invention provides for the use of antisense molecules to NABP in pharmaceutical compositions for treatment of diseases associated with the expression of NABP. The invention also describes diagnostic assays which utilize diagnostic compositions comprising the polynucleotide, fragments or the complement thereof, which hybridize with the genomic sequence or the transcript of polynucleotides encoding NABP or anti-NABP antibodies which specifically bind to NABP.

L6 ANSWER 18 OF 22 USPATFULL on STN  
AN 1999:1467 USPATFULL  
TI Human nucleic acid binding protein  
IN Bandman, Olga, Mountain View, CA, United States  
Au-Young, Janice, Berkeley, CA, United States  
Hawkins, Phillip R., Mountain View, CA, United States  
Hillman, Jennifer L., San Jose, CA, United States  
PA Incyte Pharmaceuticals, Inc., Palo Alto, CA, United States (U.S. corporation)  
PI US 5856128 19990105  
AI US 1996-698407 19960815 (8)  
DT Utility  
FS Granted  
EXNAM Primary Examiner: Wax, Robert A.; Assistant Examiner: Longton, Enrique D.  
LREP Billings, Lucy J. Incyte Pharmaceuticals, Inc.  
CLMN Number of Claims: 6  
ECL Exemplary Claim: 1  
DRWN 7 Drawing Figure(s); 7 Drawing Page(s)  
LN.CNT 1776  
CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB The present invention provides polynucleotides which identify and encode a novel human nucleic acid binding protein (NABP). The invention provides for genetically engineered expression vectors and host cells comprising the nucleic acid sequences encoding NABP. The invention also provides for the use of substantially purified NABP or its antagonists, in pharmaceutical compositions for the treatment of diseases associated with the expression of NABP. Additionally, the invention provides for the use of antisense molecules to NABP in pharmaceutical compositions for treatment of diseases associated with the expression of NABP. The invention also describes diagnostic assays which utilize diagnostic compositions comprising the polynucleotide, fragments or the complement thereof, which hybridize with the genomic sequence or the transcript of polynucleotides encoding NABP or anti-NABP antibodies which specifically bind to NABP.

L6 ANSWER 19 OF 22 USPATFULL on STN  
AN 1998:143904 USPATFULL  
TI Directed evolution of novel binding proteins  
IN Ladner, Robert Charles, Ijamsville, MD, United States  
Gutterman, Sonia Kosow, Belmont, MA, United States  
Roberts, Bruce Lindsay, Milford, MA, United States

Markland, William, Milford, MA, United States  
Ley, Arthur Charles, Newton, MA, United States  
Kent, Rachel Baribault, Boxborough, MA, United States  
PA Dyax, Corp., Cambridge, MA, United States (U.S. corporation)  
PI US 5837500 19981117  
AI US 1995-415922 19950403 (8)  
RLI Continuation of Ser. No. US 1993-9319, filed on 26 Jan 1993, now patented, Pat. No. US 5403484 which is a division of Ser. No. US 1991-664989, filed on 1 Mar 1991, now patented, Pat. No. US 5223409 which is a continuation-in-part of Ser. No. US 1990-487063, filed on 2 Mar 1990, now abandoned which is a continuation-in-part of Ser. No. US 1988-240160, filed on 2 Sep 1988, now abandoned  
DT Utility  
FS Granted  
EXNAM Primary Examiner: Ulm, John  
LREP Cooper, Iver P.  
CLMN Number of Claims: 43  
ECL Exemplary Claim: 1  
DRWN 16 Drawing Figure(s); 16 Drawing Page(s)  
LN.CNT 15973  
CAS INDEXING IS AVAILABLE FOR THIS PATENT.  
AB In order to obtain a novel binding protein against a chosen target, DNA molecules, each encoding a protein comprising one of a family of similar potential binding domains and a structural signal calling for the display of the protein on the outer surface of a chosen bacterial cell, bacterial spore or phage (genetic package) are introduced into a genetic package. The protein is expressed and the potential binding domain is displayed on the outer surface of the package. The cells or viruses bearing the binding domains which recognize the target molecule are isolated and amplified. The successful binding domains are then characterized. One or more of these successful binding domains is used as a model for the design of a new family of potential binding domains, and the process is repeated until a novel binding domain having a desired affinity for the target molecule is obtained. In one embodiment, the first family of potential binding domains is related to bovine pancreatic trypsin inhibitor, the genetic package is M13 phage, and the protein includes the outer surface transport signal of the M13 gene III protein.  
L6 ANSWER 20 OF 22 USPATFULL on STN  
AN 96:101466 USPATFULL  
TI Directed evolution of novel binding proteins  
IN Ladner, Robert C., Ijamsville, MD, United States  
Guterman, Sonia K., Belmont, MA, United States  
Roberts, Bruce L., Milford, MA, United States  
Markland, William, Milford, MA, United States  
Ley, Arthur C., Newton, MA, United States  
Kent, Rachel B., Boxborough, MA, United States  
PA Protein Engineering Corporation, Cambridge, MA, United States (U.S. corporation)  
PI US 5571698 19961105  
AI US 1993-57667 19930618 (8)  
DCD 20100629  
RLI Continuation of Ser. No. US 1991-664989, filed on 1 Mar 1991, now patented, Pat. No. US 5223409 which is a continuation-in-part of Ser. No. US 1990-487063, filed on 2 Mar 1990, now abandoned which is a continuation-in-part of Ser. No. US 1988-240160, filed on 2 Sep 1988, now abandoned  
DT Utility  
FS Granted  
EXNAM Primary Examiner: Ulm, John  
LREP Cooper, Iver P.  
CLMN Number of Claims: 83  
ECL Exemplary Claim: 1  
DRWN 16 Drawing Figure(s); 16 Drawing Page(s)

LN.CNT 15323

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB In order to obtain a novel binding protein against a chosen target, DNA molecules, each encoding a protein comprising one of a family of similar potential binding domains and a structural signal calling for the display of the protein on the outer surface of a chosen bacterial cell, bacterial spore or phage (genetic package) are introduced into a genetic package. The protein is expressed and the potential binding domain is displayed on the outer surface of the package. The cells or viruses bearing the binding domains which recognize the target molecule are isolated and amplified. The successful binding domains are then characterized. One or more of these successful binding domains is used as a model for the design of a new family of potential binding domains, and the process is repeated until a novel binding domain having a desired affinity for the target molecule is obtained. In one embodiment, the first family of potential binding domains is related to bovine pancreatic trypsin inhibitor, the genetic package is M13 phage, and the protein includes the outer surface transport signal of the M13 gene III protein.

L6 ANSWER 21 OF 22 USPATFULL on STN

AN 95:29292 USPATFULL

TI Viruses expressing chimeric binding proteins

IN Ladner, Robert C., Ijamsville, MD, United States

Guterman, Sonia K., Belmont, MA, United States

Roberts, Bruce L., Milford, MA, United States

Markland, William, Milford, MA, United States

Ley, Arthur C., Newton, MA, United States

Kent, Rachel B., Buxborough, MA, United States

PA Protein Engineering Corporation, Cambridge, MA, United States (U.S. corporation)

PI US 5403484 19950404

AI US 1993-9319 19930126 (8)

RLI Division of Ser. No. US 1991-664989, filed on 1 Mar 1991, now patented, Pat. No. US 5223409 which is a continuation-in-part of Ser. No. US 1990-487063, filed on 2 Mar 1990, now abandoned which is a continuation-in-part of Ser. No. US 1988-240160, filed on 2 Sep 1988, now abandoned

PRAI WO 1989-3731 19890901

DT Utility

FS Granted

EXNAM Primary Examiner: Hill, Jr., Robert J.; Assistant Examiner: Ulm, John D.

LREP Cooper, Iver P.

CLMN Number of Claims: 49

ECL Exemplary Claim: 1

DRWN 16 Drawing Figure(s); 16 Drawing Page(s)

LN.CNT 14368

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB In order to obtain a novel binding protein against a chosen target, DNA molecules, each encoding a protein comprising one of a family of similar potential binding domains and a structural signal calling for the display of the protein on the outer surface of a chosen bacterial cell, bacterial spore or phage (genetic package) are introduced into a genetic package. The protein is expressed and the potential binding domain is displayed on the outer surface of the package. The cells or viruses bearing the binding domains which recognize the target molecule are isolated and amplified. The successful binding domains are then characterized. One or more of these successful binding domains is used as a model for the design of a new family of potential binding domains, and the process is repeated until a novel binding domain having a desired affinity for the target molecule is obtained. In one embodiment, the first family of potential binding domains is related to bovine pancreatic trypsin inhibitor, the genetic package is M13 phage, and the protein includes the outer surface transport signal of the M13 gene III protein.

L6 ANSWER 22 OF 22 USPATFULL on STN  
AN 93:52487 USPATFULL  
TI Directed evolution of novel binding proteins  
IN Ladner, Robert C., Ijamsville, MD, United States  
Guterman, Sonia K., Belmont, MA, United States  
Roberts, Bruce L., Milford, MA, United States  
Markland, William, Milford, MA, United States  
Ley, Arthur C., Newton, MA, United States  
Kent, Rachel B., Buxborough, MA, United States  
PA Protein Engineering Corp., Cambridge, MA, United States (U.S.  
corporation)  
PI US 5223409 19930629  
AI US 1991-664989 19910301 (7)  
RLI Continuation-in-part of Ser. No. US 1990-487063, filed on 2 Mar 1990,  
now abandoned And a continuation-in-part of Ser. No. US 1988-240160,  
filed on 2 Sep 1988, now abandoned  
DT Utility  
FS Granted  
EXNAM Primary Examiner: Hill, Jr., Robert J.; Assistant Examiner: Ulm, John D.  
LREP Cooper, Iver P.  
CLMN Number of Claims: 66  
ECL Exemplary Claim: 1  
DRWN 16 Drawing Figure(s); 16 Drawing Page(s)  
LN.CNT 15410  
CAS INDEXING IS AVAILABLE FOR THIS PATENT.  
AB In order to obtain a novel binding protein against a chosen target, DNA  
molecules, each encoding a protein comprising one of a family of similar  
potential binding domains and a structural signal calling for the  
display of the protein on the outer surface of a chosen bacterial cell,  
bacterial spore or phage (genetic package) are introduced into a genetic  
package. The protein is expressed and the potential binding domain is  
displayed on the outer surface of the package. The cells or viruses  
bearing the binding domains which recognize the target molecule are  
isolated and amplified. The successful binding domains are then  
characterized. One or more of these successful binding domains is used  
as a model for the design of a new family of potential binding domains,  
and the process is repeated until a novel binding domain having a  
desired affinity for the target molecule is obtained. In one embodiment,  
the first family of potential binding domains is related to bovine  
pancreatic trypsin inhibitor, the genetic package is M13  
phage, and the protein includes the outer surface transport signal of  
the M13 gene III protein.

=>

---Logging off of STN---

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Executing the logoff script...

=> LOG Y

STN INTERNATIONAL LOGOFF AT 16:46:14 ON 05 AUG 2003